

136

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Protein

Nucleotide

Protein

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Structure

PMC

Taxonomy

OHIM

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Range: from begin to end

Reverse complemented strand

Features: SNP CDI

1: BR095794 Reports CM4-UT0076-190900...[gi:10901504] Links

LOCUS BR095794 323 bp mRNA linear EST 19-OCT-2000  
DEFINITION CM4-UT0076-190900-320-e02 UT0076 Homo sapiens cDNA, mRNA sequence.  
ACCESSION BR095794  
VERSION BR095794.1 GI:10901504  
KEYWORDS EST.  
SOURCE Homo sapiens (human)  
ORGANISM Homo sapiens  
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
Mammalia; Eutheria; Euarchontoglires; Primates; Haplorhini;  
Catarrhini; Homidae; Homo.

REFERENCE  
1 (bases 1 to 323)  
Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R.,  
Negai,M.A., da Silva,N. Jr., Zago,M.A., Bordin,S., Costa,F.F.,  
Goldman,G.H., Carvalho,A.F., Matsukuma,A., Bala,G.S., Simpson,D.H.,  
Brunstein,A., de Oliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare  
,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and  
Simpson,A.J.  
Shotgun sequencing of the human transcriptome with ORF expressed  
sequence tags  
Proc. Natl. Acad. Sci. U.S.A. 97 (7), 3491-3496 (2000)  
10737800

TITLE  
JOURNAL  
PUBMED  
COMMENT  
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This sequence was derived from the FAPESP/LICR Human Cancer Genome  
Project. This entry can be seen in the following URL  
(http://www.ludwig.org.br/scripts/gethtml2.pl?l=CM4-UT0076-190900)

FEATURES  
source  
Seq primer: puc 18 forward  
High quality sequence start: 21  
High quality sequence stop: 238.  
Location/Qualifiers  
1..323

/organism="Homo sapiens"  
/mol\_type="mRNA"  
/db\_xref="taxon:9606"  
/der\_stage="Adult"  
/clone\_lib="UT0076"  
/note="Organ: uterus; tumor; Vector: puc18; Site\_1: SmaI;  
Site\_2: SmaI; A mini-library was made by cloning products  
derived from ORESSES PCR (U.S. Letters Patent application  
No. 196,716 - Ludwig Institute for Cancer Research)  
profiles into the pUC 18 vector. Reverse transcription of  
tissue mRNA and cDNA amplification were performed under  
low stringency conditions."

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ccg gaaacacagab ttctctacaga 106  
atgagataga ggaacatatt ctactagaa 142  
acacacacac acacacacac acacacacac 181  
acacacacac acacacacac acacacacac 121  
acacacacac acacacacac acacacacac 19  
acacacacac acacacacac acacacacac 1  
ORIGIN